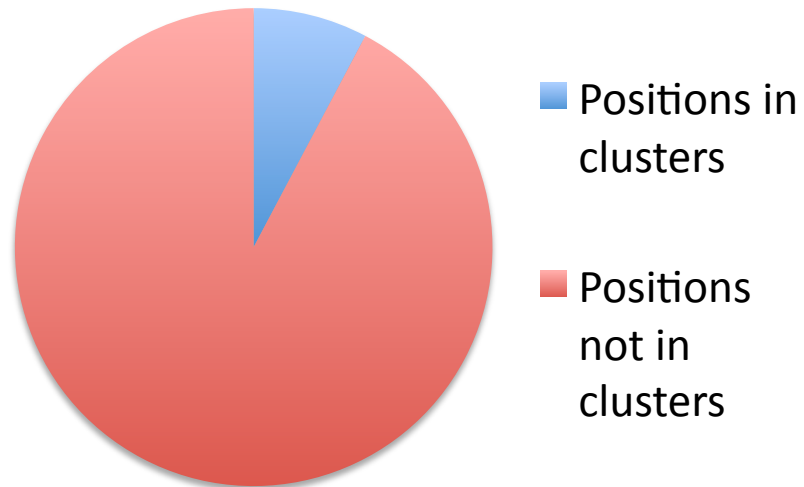
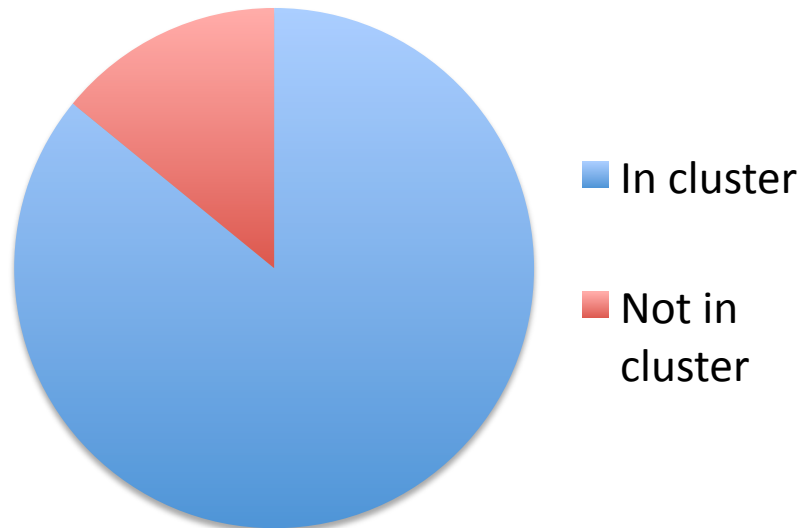


Genome Content



8.4% of positions
are in clusters

Alignment Errors



86% of alignment
errors are in clusters